# An Efficient Algorithm for Link Distance Problems (Extended Abstract)

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### **Abstract**

The link distance between two points inside a simple polygon P is defined to be the minimum number of edges required to form a polygonal path inside P that connects the points. A link furthest neighbor of a point  $p \in P$  is a point of P whose link distance is the maximum from p. The link center of P is the collection of points whose link distances to their link furthest neighbors are minimized. We present an  $O(n \log n)$ time and O(n) space algorithm for computing the link center of a simple polygon P, where n is the number of vertices of P. This improves the previous  $O(n^2)$  time and space algorithm. Our algorithm essentially sweeps a chord through the polygon and spends  $O(\log n)$  time at each step. We demonstrate that the output of the algorithm, a sequence of sets of chords, is a powerful tool for solving several other link distance problems.

## 1 Motivations and Definitions

The study of link distance problems was motivated by the following robot motion-planning problem. Consider a point-size robot who wants to move between two points within a polygon P. Suppose the robot can perform two types of movements: a straight line motion and a pure rotation. Therefore a feasible path for the robot is a polygonal chain that connects the two points in P. Further suppose that straight line motion is cheap, while a pure rotation is computationally expensive. Therefore a "good" path for the robot is the one that has as few number of turns, or number of edges (links), as possible.

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We now give several formal definitions concerning the link distance between points. A minimum link path between two points  $p_1$  and  $p_2$  is a polygonal path that connects  $p_1$  and  $p_2$  in P with the minimum number of edges; this number is the link distance between  $p_1$  and  $p_2$ . The problem of computing a minimum link path was solved by Suri [12], in which an  $O(n \log \log n)$ time algorithm, or O(n) time if a triangulation of P is precomputed [11][14], was presented, where n is the total number of vertices of P. Given a point  $p \in P$ , its link furthest neighbor is a point of P whose link distance from p is the maximum over all other points of P. The link center of P is the collection of points whose link distances to their link furthest neighbors are minimum over all other points of P. The link radius of P is the link distance from a point in the link center to its link furthest neighbor. The link diameter of P is the maximum link distance between two points of P, or equivalently, the maximum link distance from a point to its link furthest neighbor over all points of  $\boldsymbol{P}.$ 

We now list some of our results and compare them with the best previous ones.

- Computing the link center and radius in  $O(n \log n)$  time and O(n) space (best previous result:  $O(n^2)$  time and  $O(n^2)$  space  $|10|^{-1}$ );
- Computing the link diameter in  $O(n \log n)$  time and O(n) space (same complexities [13]);
- Solving the all link furthest neighbor problem in  $O(n \log n)$  time, that is, report a link furthest neighbor for each vertex of P (best previous result:  $O(n^2)$  time and O(n) space [13]);
- Preprocessing P in  $O(n \log n)$  time and O(n) space such that given a query point in P, its link furthest neighbor and the link distance between

<sup>&</sup>lt;sup>1</sup>As this abstract was been written, we learned of [2], which presents an  $O(n \log n)$  time and O(n) space algorithm to compute a non-empty region within the link center. Later they claimed that the algorithm was improved to find the entire link center.

them can be found in  $O(\log n)$  time (best previous result:  $O(n \log \log n)$  time and O(n) space preprocessing, and O(n) query time [13]);

Besides these, our algorithm can also solve some extensions of the link distance problems. First we need more definitions. A minimum link path between a point p and a line segment s is a polygonal path that connects p to a point on s within P with the minimum number of edges; this number is the link distance between p and s. A link furthest neighbor of a line segment  $s \in P$  is a point of P whose link distance from s is the maximum over all other points of P. A link central segment of P is a line segment s whose link distance to its link furthest neighbors is the minimum over all other segments in P;

In the following, we list our results for two problems that, as far as we know, have not been studied previously.

- Computing a shortest link central segment in  $O(n \log n)$  time and O(n) space;
- Preprocessing P in O(n log n) time and O(n) space such that given a query segment, its link furthest neighbor and the link distance between them can be found in O(log n) time;

All these results are achieved essentially by one algorithm, which uses a *chord* to sweep through the polygon P in a recursive fashion. It generates a sequence of sets of chords that satisfy a certain *Intersection Properties*, Property I and  $I^*$ , which provides all the information we need to solve the mentioned problems.

In section 2, we introduce the concept of Property I and  $I^*$ , and show how to solve all the link distance problems by using the sets of chords with this property. Then in Section 3 and 4, two algorithms are described that generate the desired sets of chords: a brute-force algorithm and a simple  $O(n^2 \log n)$  sweep-chord algorithm. Section 5 improves the time complexity to  $O(n \log n)$ . Finally in Section 6, we show our results for the additional extensions of link distance problems.

# 2 Intersection Property

We start with some basic definitions. A chord of P is a line segment in the interior of P except for its two endpoints, which then must be on the boundary of P. Therefore a chord c divides P into exactly two subpolygons, and one of them (specified later) is called the *component* of c, denoted by P(c).

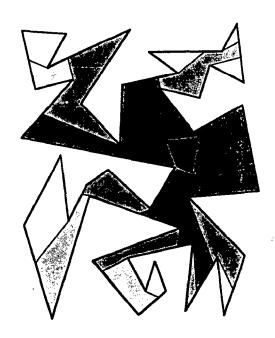


Figure 1: The partition of P

A sequence of sets of chords  $C_1, C_2, ..., C_k$  is said to have the *Intersection Property*, or Property I, if the following statement holds:

For any point  $p \in P$ , p has at most link distance i,  $1 \le i \le k$ , to every point of P iff it is contained in the intersection region  $R_i = \bigcap_{c \in C_i} P(c)$ .

Suppose such  $C_i$  is available,  $1 \le i \le k$ . Let us see how can we use them to solve the previously mentioned link distance problems. First, since any point  $p \in P$  can reach every point of P by at most d links, where d is the link diameter of P, we immediately conclude that d = k + 1. Clearly the link center is  $R_r$ , where r is the link radius and satisfies the inequality  $\lceil d/2 \rceil \le r \le \lceil d/2 \rceil + 1 \lceil 10 \rceil$ . Therefore the link center and radius can be computed in  $O(N \log N)$ time, where N is the total number of chords in the sets. (It can be shown that  $N = \Omega(n)$ .) Next define  $R'_r = R_r, R'_i = R_i - R_{i-1}$  for  $r+1 \le i \le d-1$ , and  $R'_{d} = P - R_{d-1}$ . Then any points in  $R'_{i}$  can reach all points of P by i links, but not i-1 links. In other words, they have link furthest neighbors that are i links away. These regions  $R'_r$ ,  $R'_{r+1}$ , ...,  $R'_{d-1}$ ,  $R'_d$  form a partition of P. See Figure 1 for an example: the black region is the collection of points having link distance 4 to their link furthest neighbors; this region is the link center and 4 is the link radius of the polygon. The dark shaded region is the collection of points having link distance 5 to their link furthest neighbors. The shaded region is the collection of points having distance 6 to their furthest neighbors. The light shaded region having distance 7 to their neighbors. The blank region having distance 8 to their neighbors, 8 is the link diameter of the polygon.

Using the point-location techniques [3], we can preprocess this partition in  $O(N \log N)$  time and O(N) space such that given a query point p, its link furthest neighbor and the link distance between them can be obtained in  $O(\log N)$  time.

A stronger version of the Intersection Property, Property  $I^*$ , is defined as follows:

For any line segment  $s \in P$ , s has at most link distance i,  $1 \le i \le d-1$ , to every point of P iff it intersects the component P(c) for each chord  $c \in C_i$ .

We use the following results from [8]: given a chord set C with size k in a n-gon P, we can decide in  $O(n \log n + k \log k)$  time and O(n + k) space, or  $O(\log n + \log k)$  time if an  $O(n \log n + k \log k)$  time and O(n+k) space preprocessing of C and P is allowed, whether there exists a line segment intersecting the component of each chord in C; if the answer is positive, then we can find a shortest one in  $O(n \log n + k \log k)$ time and O(n+k) space. Therefore computing a shortest link central segment costs  $O(N \log N)$  time once we have the chord set  $C_i$ ,  $1 \le i \le d-1$ . We can also solve the query problem as follows: first preprocess  $C_i$ , for  $1 \le i \le d-1$ , by the algorithm in [8]. Then when a query segment s is given, choose an arbitrary point  $p \in s$  and find the link distance, say i, between p and its link furthest neighbor; then the link distance from s to its link furthest neighbor is either i or i-1, which can be found out in additional  $O(\log N)$  time.

Given the power of sequence of chords satisfying Property I and  $I^*$ , there are two goals: to find sets of total size N = O(n), and to do so quickly, in  $O(n \log n)$  time.

# 3 A Brute-force Algorithm

In this section, we briefly describe a brute-force algorithm that generates the desired chord sets  $C_1, C_2, ..., C_{d-1}$  with  $O(n^2)$  total number of chords. For simplicity, we ignore all degeneracies throughout the paper. It is well known that any point  $p \in P$  can reach all points of P by i links iff it can reach all vertices of

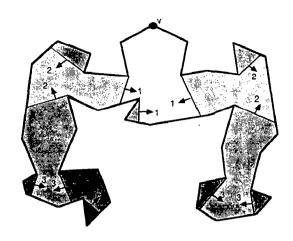


Figure 2: An example of the sets of chords with respect to a vertex v. The arrows on the chords indicate at what sides of their components lie. The number at each chord indicates the index of the set that contains that chord.

P by i links. we will use this fact implicitly in many places.

We start with one vertex v, and construct its visibility polygon V(v). Let  $C_1(v)$  be the set of chords on the boundary of V(v). For each  $c \in C_1(v)$ , define its component P(c) to be the sub-polygon that contains v. Then  $\bigcap_{c \in C_1(v)} P(c) = V(v)$  is exactly the collection of points that can reach v by at most 1 link. We remove V(v) from P and, within its remaining portion, construct the visibility polygon V(c) for each chord  $c \in C_1(v)$ . Let  $C_2(v)$  be the set of newly generated chords with their components defined in the same manner. Then  $\bigcap_{c \in C_2(v)} P(c)$  is exactly the collection of points that can reach v by at most 2 links. We can repeat this construction until the entire polygon P is covered. In the end, we have a sequence of chord sets  $C_1(v)$ ,  $C_2(v)$ , ...,  $C_{d(v)-1}(v)$ , where d(v) is obviously the link distance from v to its link furthest neighbor. See Figure 2. The entire procedure can be managed to run O(n) time provided that a triangulation of Pis given (see [13] for details).

At this time, We need to give a few definitions for our later discussion. We define the *label* of a chord  $c \in C_i(v)$  to be  $i, 1 \le i \le d(v) - 1$ . A chord c is called a *direct descendant* of another chord c' if c is on the boundary of V(c'), i.e., c is generated from c'; and c' is called the *direct ancestor* of c. We say that  $c_k$  is a descendant of  $c_1$  if there is sequence of chords  $c_k$ ,  $c_{k-1}$ , ...,  $c_2$ ,  $c_1$  such that  $c_i$  is a direct descendant of

 $c_{i-1}$ , for i=k,k-1,...,3,2; and  $c_1$  is an ancestor of  $c_k$ . There is a simple observation: each chord c has one endpoint incident to a reflex vertex of P, which we call the *origin* of c; the other endpoint we call the *hit point* of c. One can understand this definition by imagining the direct ancestor of c (each chord has a unique direct ancestor) to be a light source; then c is a light ray shooting from its origin to its hit point.

We repeat the above described procedure for each vertex of P. Let  $d = \max_{\text{each vertex } v} d(v)$  to obtain the link diameter of P. Then let  $C_i = \bigcup_{\text{each vertex } v} C_i(v)$ , for  $1 \le i \le d-1$ , to obtain the sets of chords that have Property I and  $I^*$ . The total number of chords is clearly  $O(n^2)$ .

# 4 A Sweep-chord Algorithm

The improvement over the brute-force algorithm is possible only if we can bound the total number of chords in the sets. Consider a chord  $c \in C_i$ . We define c to be redundant if there exists another chord  $c' \in C_i$  such that  $P(c') \subset P(c)$ . Intuitively, if c is redundant, then P(c) has no contribution to the intersection  $\bigcap_{c \in C_i} P(c)$ . A chord is necessary if it is not redundant. There are immediately two observations. First, the two chords c and c' need not have the same label. c is also redundant when the label of c' is greater than that of c. Second, all descendants of a redundant chord are also redundant.

We now show that there are only O(n) necessary chords in the sets  $C_1, ..., C_{d-1}$ . Recall that each chord c has one endpoint incident to a reflex vertex v of P. We define c to be a clockwise chord if its component lies in clockwise direction about v. Similarly, define c to be counterclockwise if its component lies in counterclockwise direction about v. We next show that there are at most two necessary chords incident to v, one is clockwise and the other counterclockwise. Suppose there are two, say clockwise, chords  $c_i$  and  $c_j$  incident to v. If they have the same label, then one of them is redundant. So wlog, let i and j be their labels respectively such that i < j. Then we have two cases. If  $c_i$  lies in clockwise direction from  $c_i$ , then  $c_i$  is redundant. Now consider the case when  $c_i$  lies in clockwise direction from  $c_j$ . Consider the direct ancestor  $c'_j$  of  $c_i$ . We know  $c'_i$  is contained in  $P(c_i)$ . Suppose it is not contained in  $P(c_i)$ . Then  $c'_i$  must intersect  $c_i$  at a point p that can see a small neighborhood of v, including some points not in  $P(c_i)$ . This is not possible. Therefore  $c'_i$  is contained in  $P(c_i)$ , implying that  $P(c'_i)$ is contained in  $P(c_i)$ . Since j > i, the label of  $c'_j$  is no less than i. So  $c_i$  is redundant. We thus conclude that there are O(n) necessary chords.

Our algorithm generates O(n) number of chords that include all the necessary ones. We give one definition first. We say a chord c is downward if its origin is above its hitpoint (again, imagine the direct ancestor of c to be a light source, then c is a ray shooting from its origin down to its hit point); otherwise c is called upward.

As a preprocessing step of the algorithm, we first partition P into monotone regions as follows. For each reflex vertex v that either points downwards (we call it a downward cusp) or upwards (upward cusp), we draw the two horizontal chords that are incident to v. This gives us a monotone partition. Then we define a graph whose nodes correspond to the monotone regions and two nodes are connected by an edge if their corresponding regions are adjacent. We will occasionally use the same notation for a node and its corresponding region. The defined graph is clearly an O(n) size tree. Let T be the obtained embedding of the tree. Within T, if a node  $t_1$  is adjacent to another node  $t_2$ from above, then  $t_1$  is called a parent of  $t_2$ , and  $t_2$  a child of  $t_1$ . Two nodes are called siblings (spouses) if they have a common parent (child). A top node is the one that has no parents, and a bottom node has no children.

Unlike the brute-force algorithm that generates all the descendants of each vertex, one at a time, the sweep-chord algorithm basically takes care of all downward (or upward) chords incident to each reflex vertex, one at a time, by a recursive sweep of the entire polygon. In this way, the generation of redundant chords is well under control. The general structure of the algorithm is as follows.

#### ALGORITHM SWEEP-POLYGON

- 1. Let all nodes of T be unmarked;
- 2. for each bottom node t of T do
- 3. SWEEP-DOWN(t);
- 4. for each top node t of T do
- 5. SWEEP-UP(t);

END of SWEEP-POLYGON.

There are two procedures in the algorithm. The procedure SWEEP-DOWN (t) basically sweeps the monotone region t downwards, and then marks the node t. The procedure SWEEP-UP (t) operates similarly. Since the two procedures are symmetric, we only describe the former one.

#### **PROCEDURE** SWEEP-DOWN (t)

1. if t has two parents  $t_1$  and  $t_2$  then

```
2.
       for i = 1 to 2 do
3.
           if t_i is not marked by "d" then
4.
              SWEEP-DOWN(t_i);
    else if t has one parent t_1 and one sibling
5.
    t_2 then
6.
       if t_1 is not marked by "d" then
7.
           SWEEP-DOWN(t_1);
8.
        if t_2 is not marked by "u" then
9.
           SWEEP-UP (t_2);
10.
    Sweep the monotone region t downwards
    to generate all necessary downward chords
    within t, then mark t by "d";
END of SWEEP-DOWN(t).
```

A simple observation is that each monotone region is processed downwards and upwards, each exactly once. Within each monotone region, the algorithm sweeps a horizontal chord  $c_s$  from top to bottom and stops at each vertex to update the maintained data structure and generate downward chords incident to that vertex. Upon the completion of the algorithm, all necessary chords should be generated. More precisely, Step 10 consists of the following three stages:

- Initial stage: Collect the information obtained from  $t_1$  and  $t_2$ . Then build up the appropriate data structure;
- Sweeping stage: Sweep the monotone region downwards vertex by vertex, and maintain the data structure;
- Processing stage: When the entire monotone region has been swept, process the data structure such that they can be used by the sweeping procedure of other monotone regions;

We now sketch the data structure maintained by the algorithm. Let  $c_s$  be the current sweeping chord in a downward sweep. To keep track of all the downward chords below  $c_s$ , we have to maintain all their ancestors above  $c_s$ . More precisely, we call a chord c active with respect to  $c_s$  if (1) it is above  $c_s$ , (2) its component is disjoint from  $c_s$ , and (3) it is visible from  $c_s$ . Intuitively, these are the chords that have downward direct descendants below  $c_s$  to be constructed. It is sufficient and necessary for the algorithm to maintain all the active chords with respect to  $c_s$  to generate the downward chords below  $c_s$ . However, we can not afford to maintain all the active chords separately since there are O(n) of them.

The solution is to organize the O(n) number of active chords into two convex chains, which can be more easily manipulated geometrically. First we throw away

all the chords whose labels are less than M-1, where M is the maximum label of the active chords. Then we obtain a partition of all active chords with label M (the treatment for the chords with label M-1is identical and thus omitted). Within each class of the partition, we take the intersection of the components of all the chords; the boundary of the intersection region in the interior of P is a convex chain. The partition is chosen in such a way that all intersection regions are disjoint. Therefore all convex chains are disjoint and naturally ordered along the boundary of P. It is shown that these convex chains, together with those obtained from active chords with label M-1, will give all the necessary downward descendants below c<sub>a</sub>. Furthermore, every non- (left or right) extreme convex chain can be replaced by its direct descendants that cross the sweeping chord  $c_s$ ; we call these direct descendants critical. Therefore to generate all necessary downward descendants, the algorithm needs to know the four convex chains (the leftmost and rightmost with label M, and with label M-1; these are called the active chains, or ACs for short), and all critical chords. We called this the upper data structure of cs, or UD for simplicity. The lower data structure LD can be similarly defined.

We assume that P has been preprocessed for bullet shooting and geodesic path queries. The first one is to find the hit point of a query ray from a given a point in a given direction. An O(n) time algorithm, provided that P is triangulated, was given by Chazelle and Guibas that can build an O(n) space data structure with  $O(\log n)$  query time [1][5]. The second one is to output the geodesic path between two query points. For this, an O(n) space data structure was proposed by Guibas and Hershberger [4]. It can be constructed in O(n) time, provided that P is triangulated, and performs the query in  $O(\log n)$  time, under the condition that the output path will be returned before the next query is made. These two data structures will be extensively used by our algorithm.

As mentioned above, the algorithm performs exactly two sweeps for each node of T, one downward and one upward. Let t be a node of T and  $R_t$  be its corresponding monotone region. We use  $c_{tt}(t)$  and  $c_t(t)$  to denote the upper and lower horizontal boundaries of  $R_t$  (each of them can either be a chord, or consists of two chords). We will only describe the downward sweep of t.

#### 1. Initial stage: We have three cases:

a. t is a top node of T. We create an empty UD with respect to  $c_u(t)$ , with ACs assigned the la-

bel 0.

- b. t has two parents t<sub>1</sub> and t<sub>r</sub> with t<sub>1</sub> to the left of  $t_r$ . Then  $c_u(t)$  consists of two chords  $c_l(t_l)$  and  $c_l(t_r)$ . By the algorithm, both  $t_l$  and  $t_r$  have been processed downwards. Therefore both  $UD_l$ and  $UD_r$ , which are the UDs with respect to  $c_l(t_l)$  and  $c_l(t_r)$ , are available. We obtain the UD with respect to  $c_u(t)$  by merging  $UD_l$  and  $UD_r$ . More precisely, let  $k_l$  and  $k_l-1$  be the labels of ACs in  $UD_l$ ,  $k_r$  and  $k_r - 1$  the labels of ACs in  $UD_r$ . Wlog, let  $k_l \leq k_r$ . If  $k_l =$  $k_r = k$ , then the left extreme AC in  $UD_l$  and right extreme AC in  $UD_r$ , with labels k and k-1, form the ACs in UD. If  $k_1=k_r-1$ , then the ACs in  $UD_l$  with label  $k_l - 1$  will be discarded. The left extreme AC with label  $k_l$  in  $UD_l$  and the right extreme AC with label  $k_r-1$ in  $UD_r$ , plus the ACs with label  $k_r$  in  $UD_r$ , form the ACs in UD. If  $k_l \leq k_r - 2$ , then ACs in  $UD_l$  will completely discarded. the ACs in  $UD_r$ will be the ACs in UD. In addition, the set of critical chords are combined correspondingly in each case.
- c. t has one parent tp and one sibling ts. Then  $c_l(t_p)$  consists of  $c_u(t)$  and  $c_u(t_s)$ . By the algorithm,  $t_p$  ( $t_s$ ) has been processed downwards (upwards). Therefore both  $UD_p$  and  $LD_s$ , which are the UD and LD with respect to  $c_l(t_p)$ and  $c_u(t_s)$  respectively, are available. We obtain the UD with respect to  $c_u(t)$  by merging  $UD_p$  and  $UD_s$ . Here we describe some details. Wlog, let  $t_s$  be to the left of t. Notice some critical chords in DL, may be active with respect to  $c_u(t)$ . Consider those of them with the maximum label  $k_s$  (incidentally, these are upward chords). Let c be the one whose hit point is ordered first clockwise along the boundary of P starting at the left endpoint of  $c_u(t_s)$ . Let  $k_p$ and  $k_p - 1$  be the labels of ACs in  $UD_p$ . Then we have several cases to consider. If  $k_s \leq k_p - 1$ , then UD is the same as  $UD_p$ . If  $k_s = k_p$ , then ACs in UD includes the AC's in  $DU_p$  with label  $k_p - 1$ , plus the modified ACs (by c) in  $DU_p$ with label  $k_p = k_s$ . If  $k_s = k_p + 1$ , then ACs in UD includes c, and the ACs in UD with label  $k_p$ . Finally, if  $k_s \geq k_p + 2$ , then c is the only AC in UD and the entire  $UD_p$  is discarded. In addition, the set of critical chords are computed correspondingly in each case.

The important observation is that all ACs discarded in the procedure will produce only redundant chords. Therefore their absence does not effect the correctness of the final output. The critical chord c found in Case c essentially takes care of the situation when a link path travels out the monotone region  $t_s$ , makes a turn, and then goes into the monotone region t. We thus call c the T-chord (T stands for turn). It should be mentioned that the T-chord c is not computed in this stage, it was actually found and stored at  $c_u(t_s)$  in the processing stage of  $t_s$  (which will be described later). Therefore an initial stage of the algorithm can be implemented in  $O(\log n)$  time.

2. Sweeping stage: Each time the sweeping chord  $c_s$  stops at a vertex, we check if any critical chords have become active (that is, above  $c_s$ ). Then these chords are used to update the current ACs and generate new critical chords. Details can be found in [6].

Within each step of the sweeping stage, it costs O(n) time to find those out-of-date critical chords, and  $O(\log n)$  time to update the current ACs. The new ACs must be able to generate all necessary descendants that the old ACs have.

- 3. Processing stage: Again, we have three cases for t:
  - a. t is a bottom node. Then the processing stage is
  - b. t has a spouse  $t_s$  and a child  $t_c$ . Then  $c_u(t_s)$  consists of  $c_l(t)$  and  $c_l(t_s)$ . Wlog, let t be to the left of  $t_s$ . Our task in this case is to find the critical chord c in UD with the maximum label, and whose hit point is ordered last (among all critical chords with the maximum label) clockwise along the boundary of P starting from the left endpoint of  $c_l(t)$ . The chord c is the T-chord with respect to  $c_l(t_s)$ .
  - c. t has two children  $t_l$  and  $t_r$ . Then  $c_l(t)$  consists of  $c_u(t_l)$  and  $c_u(t_r)$ . Our task in this case is to compute the UDs with respect to  $c_u(t_l)$  and  $c_u(t_r)$  by splitting UD with respect to  $c_l(t)$ . Notice some critical chords may be active with respect to  $c_u(t_l)$ , and some be active with respect to  $c_u(t_r)$ . Using a similar criterion for choosing T-chord in Case c of initial stage, we choose one of them for  $c_u(t_l)$ , and one for  $c_u(t_r)$ . These two chords are called S-chords (S stands for straight) The two S-chords will then be used to obtain UDs with respect to  $c_u(t_l)$  and  $c_u(t_r)$ .

A processing stage can be implemented in  $O(n \log n)$  time. The algorithm thus takes overall  $O(n^2 \log n)$  time. The correctness of the algorithm is based on the fact that at each step, we only discard ACs and critical chords that have no contributions to the necessary chords.

## 5 The Improvement

The the bottleneck for a more efficient algorithm is the  $O(n \log n)$  time complexity for a processing stage and the O(n) time complexity for a step in a sweeping stage. These are the two issues we will deal with in this section.

The algorithm can generate a critical chord in one of the three stages. Suppose at some moment, the algorithm generates a downward critical chord c. We observe that c will not be used by the algorithm until one of the following events occurs:

- In a sweeping stage, c becomes active with respec to the sweeping chord c, as it moves down;
- In a processing stage, c crosses the sweeping chord
  c, that is incident to a downward cusp. At this
  moment, c may become a T-chord;
- In a processing stage, c crosses the sweeping chord
  c, that is incident to an upward cusp. At this
  moment, c may become a S-chord;

The idea behind the improved algorithm is as follows. We will not main the critical chords in UD (and LD). In stead, we insert a critical chord c, as soon as it is created, into some data structures such that c can be captured later in the algorithm when it is needed. There are two such data structures. The first one is used for the occurrence of the first event. It consists of a collection of convex chains, each for an edge of P. When a critical chord c is created, we first find the edge of P that contains the hit point of c, then update the associated convex chain appropriately. The second data structure consists of two balanced trees  $T_d$  and  $T_u$ that are used for the second and third events. In the remainder of the section, we will very briefly describe  $T_u$  ( $T_d$  is similar) and the idea of how it improves our algorithm.

The leaves of  $T_u$  are all the upward cusps of the polygon P, and they are arranged according to a linear order defined below after a few definitions. Let v be an upward cusp of P and s be the union of two horizontal chords incident to v. (We call the one to the left the L-chord, and the other one the R-chord; these two chords are also called paired). The polygon P is divided by s into three components. The one above s is called the upper component of v. For the two remaining components below s, the one to the left of v in its small neighborhood is called the left component of v, and the other one the right the right component of v. An upward cusp is called a top upward cusp if its upper component contains no other upward cusps.

Choose an arbitrary top upward cusp v. We define all the upward cusps in its left component to be ordered lower than v, and all those in its right component are ordered higher than v. Then the definition is applied recursively to a top upward cusp in each of its left and right components. All L-chords (R-chords) can also be ordered by their incident upward cusps. Therefore we can also think of leaves of  $T_u$  as all L-chords (R-chords). Each internal node of  $T_u$  is associated with the following four fields: LT, LS, RT and RS (they stand for left T-chord, left S-chord, right T-chord and right S-chord respectively).

In a moment, we will show that this order can be computed efficiently in  $O(n \log n)$  time. First let us see how  $T_u$  is used in the improved the algorithm. Suppose a new critical chord c is created at a step of the algorithm. Wlog, let c be a downward critical chord with its component at the left side (for this, c is called the left critical chord). Then c is a potential S-chord of some R-chords if c intersects their paired L-chords. So consider all the L-chords that are intersected by c. Let  $c_a$  and  $c_b$  be the highest and lowest among them along c. In fact, they are the highest and lowest ordered ones among such L-chords. For this reason, they are called the *limiting* chords of c. We store c into the LS fields of a subset of nodes in  $T_u$  whose leaves exactly cover all such L-chords (this set of nodes has size  $O(\log n)$  and can be found in  $O(\log n)$  time). In a similar way, c will also be inserted into the fields LT of an appropriate subset of nodes in  $T_d$ . In general, we have the following rules:

- 1. c is downward left: insert into LS fields of  $T_u$  and LT fields of  $T_d$ ;
- 2. c is upward left: insert into LT fields of  $T_u$  and LS fields of  $T_d$ ;
- 3. c is downward right: insert into RS fields of  $T_u$  and RT fields of  $T_d$ ;
- 4. c is upward left: insert into RT fields of  $T_u$  and RS fields of  $T_d$ ;

By the time we want to compute, e.g., a S-chord for a R-chord, we simply locate its paired leaf in  $T_u$  and walk up to the root to examine the chords stored in the LS fields of its ancestors. Unfortunately, there could be total O(n) chords stored on the path by the naive approach. However, we show next that at most one chord needs to be stored at each node of  $T_u$ .

Suppose a downward left critical chord  $c_2$  is being inserted into the LS field of a node t in  $T_u$  when we discover that another chord  $c_1$  is already there. Then

we first compare their labels and discard the one with less label. When there is a tie, we further check the positions of their components. Suppose they are disjoint. Consider their higher limiting chords and the corresponding upward cusps. We discard one of  $c_1$  and  $c_2$  that is contained in the upper component of the upward cusp of the other (it can be shown that such a chord always exists). Now suppose the components of  $c_1$  and  $c_2$  are not disjoint. Then we discard the one whose origin is outside the component of the other.

Wlog, let  $c_2$  be discarded. We have to show that  $c_2$  can not be the necessary S-chord for a R-chord  $c_r$ , where  $c_r$  is also a leaf descendant of t. Let  $c_l$  be the paired chord of  $c_r$ . They are both incident to an upward cusp v. There are several cases to consider.

- I.  $c_2$  does not intersect  $c_l$ . In this case,  $c_2$  can not be the S-chord of  $c_r$  by definition.
- II.  $c_2$  intersects  $c_l$ . Then  $c_2$  is a potential S-chord of  $c_r$ . We further have two cases.
  - A.  $c_1$  intersects  $c_l$ . We know that  $c_1$  is the winner. This implies that either the label of  $c_1$  is greater than that of  $c_2$ , in which case  $c_2$  can not be the S-chord of  $c_r$  (similar to the argument in Case c of the processing stage), or the origin of  $c_1$  is contained in the component  $P(c_2)$  of  $c_2$ . An equivalent way to say this is that the origin of  $c_1$  is ahead of that of  $c_2$  clockwise along the boundary of P starting at the left endpoint of  $c_l$ . Therefore by the same argument,  $c_2$  is redundant.
  - B.  $c_1$  does not intersect  $c_l$ . Let  $c_a$  and  $c_b$  be the limiting chords of  $c_1$  with their incident upward cusps  $v_a$  and  $v_b$ . We claim that there are the following two possibilities: 1. the left component of v contains the hit point of  $c_1$ ; 2. there exists an upward cusp  $v_a < v' < v$  whose left component contains the hip point of  $c_1$ , and whose right component contains v (Actually, the correctness of this claim does not depend on the fact that  $c_1$  does not intersect  $c_l$ ). We omit the proof of the claim in this paper.
    - a. The left component of v contains the hit point of c<sub>1</sub>. We know that c<sub>1</sub> does not intersect c<sub>l</sub>. Therefore c<sub>1</sub> is completely contained in the left component of v. But then v would not be included by the limiting chords c<sub>n</sub> and c<sub>h</sub>, and therefore c would not be inserted into the LS field of t. Thus this case is not pos-

sible.

- b. There exists an upward cusp  $v_a < v' < v$  whose left component contains the hip point of  $c_1$ , and whose right component contains v. Let  $c'_l$  and  $c'_r$  be the paired chords incident to v'. Clearly,  $c_1$  can not be completely contained in the left component of v'. Therefore  $c_1$  must intersect  $c'_l$ . We have two cases.
  - i. The labels of  $c_1$  and  $c_2$  are the same. Since  $c_1$  is the winner,  $c_1$  and  $c_2$  must be disjoint but not their components. Therefore  $P(c_2)$  is contained in  $P(c_2)$ , implying that  $c_2$  is redundant.
  - ii. The label of  $c_1$  is greater than that of  $c_2$ . Then any direct descendants of  $c_2$  below  $c_r$  have their components containing that of  $c_1$  but with no greater label. Therefore  $c_2$  is not the necessary S-chord of  $c_r$ .

Therefore we only have to look at  $O(\log n)$  chords on the path. We conclude that an insertion and extraction can be done in  $O(\log n)$  time.

We devote the remaining section to the description of an  $O(n \log n)$  algorithm that computes the order defined on upward cusps. First we partition P with all L- and R-chords. Each upward cusp can be uniquely associated with a region in the partition directly above it. We then define a graph G as follows. The nodes of G are all upward cusps, and an edge is added between two nodes iff their associated regions are adjacent in the partition. The graph G is a tree of size O(n), and can be constructed in  $O(n \log n)$  time. Following the definitions in Section 2, it is clear that an upward cusp is a top node in G iff it is a top upward cusp. We can find the set S of all top nodes of G in O(n) time. We assume that all nodes of S are stored in a linked list so that they can be accessed in constant time by their indices. The output of our  $O(n \log n)$  algorithm is a rooted binary tree T' such that the nodes of T' are all the upward cusps, and the inorder of them are exactly the order defined above. To initialize the algorithm, we choose an arbitrary top node t of G, and let it be the root of T'. Then t is deleted from G together with its two incident edges. This will split G into two components  $G_l$  and  $G_r$  such that  $G_l$  contains the left child  $t_l$  of t, and  $G_r$  contains the right child  $t_r$  of t. Let  $S_l$  and  $S_r$  be the set of top nodes of  $G_l$  and  $G_r$ respectively. Algorithmically, the two sets  $S_l$  and  $S_r$ can be obtained as follows. We choose the one of  $G_l$ and  $G_r$ , say  $G_l$ , that has fewer nodes than the other. We can accomplish this selection in  $O(|G_l|)$  time by

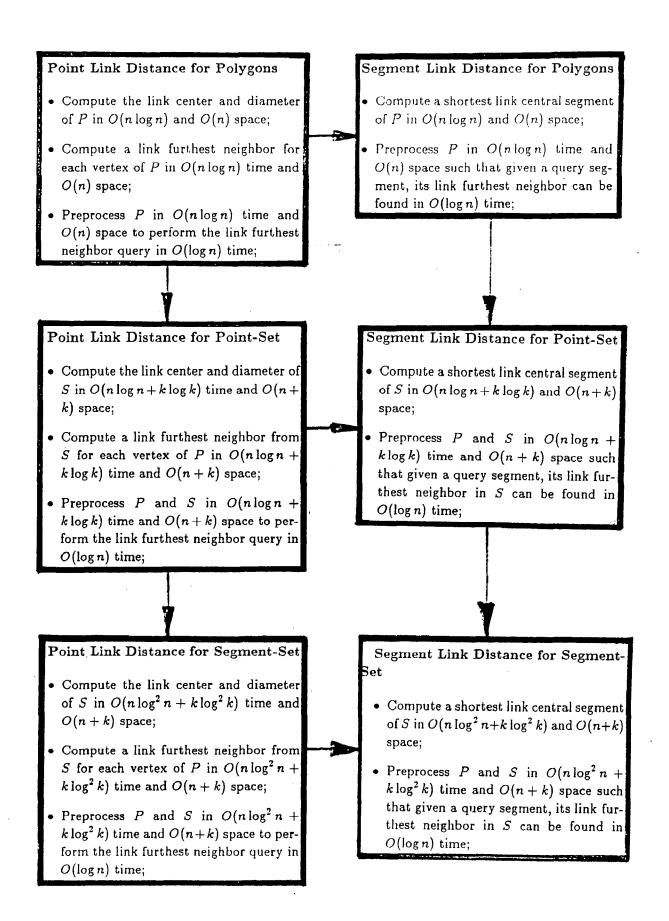


Figure 3. The table of results. The arrows indicate the directions of generalizations.

counting the nodes of  $G_l$  and  $G_r$  alternatively until one of them is exhausted. Then in additional  $O(|G_l|)$  time, we can find the set  $S_l$ . Then all nodes of  $S_l$ , except possibly  $t_l$ , are deleted from  $S_u$ . The remaining nodes in  $S_u$ , possibly plus  $t_r$ , form the set  $S_r$ . After this, the algorithm can recursively choose an arbitrary node from  $S_l$  to be the left child of t in  $T'_u$ , and a node from  $S_r$  to be the right child of t in  $T'_u$ , and so on. The time complexity of the algorithm is  $O(n \log n)$ .

## 6 Conclusion

Several extensions of the link distance problem can be solved by the described algorithm very efficiently. Let S be a set of k points (or line segments) in P. We define the link center of S to be the collection of points of P whose maximum link distance to points (or line segments) in S is the minimum, and the link diameter of S to be the maximum link distance achieved by two points (or line segments) in S. We assume that points (or line segments) in S are not obstacles for the link paths; only the boundary of P is. In general, we can redefine all the previous problems with respect to the set S (instead of the points of P). Using the sweep-chord algorithm and the techniques in [9], we can achieve the results listed in Figure 3 [7].

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